

NCBI Nucleotide

Entrez PubMed Nucleotide Protein Genome Structure PMC Taxonomy

Search Nucleotide for glucosylceramide synthase

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1: Reports Links
NM_078456

Caenorhabditis elegans ceramide glucosyltransferase / glucosylceramide synthase family member (50.1 kD) (XS315) mRNA
 gi|25147525|ref|NM_078456.2|[25147525]

Related resources **2:** Reports Links

NM_127546

Arabidopsis thaliana ceramide glucosyltransferase, putative (At2g19880) mRNA, complete cds
 gi|18399183|ref|NM_127546.1|[18399183]

3: Reports Links
XM_538788

PREDICTED: Canis familiaris similar to Ceramide glucosyltransferase (Glucosylceramide synthase) (GCS) (UDP-glucose:N-acylsphingosine D-glucosyltransferase) (UDP-glucose ceramide glucosyltransferase) (GLCT-1) (LOC481666), mRNA
 gi|57094232|ref|XM_538788.1|[57094232]

4: Reports Links
NW_139871

Canis familiaris chromosome 11 genomic contig, whole genome shotgun sequence
 gi|54126007|ref|NW_139871.1|Cfa11_WGA36_1|[54126007]

5: Reports Links
NM_031795

Rattus norvegicus UDP-glucose ceramide glucosyltransferase (Ugcg), mRNA
 gi|25742622|ref|NM_031795.1|[25742622]

6: Reports Links
NM_004775

Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase,

polypeptide 6 (B4GALT6), mRNA
gi|13929471|ref|NM_004775.2|[13929471]

7: Reports Links
NM_003358

Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA
gi|4507810|ref|NM_003358.1|[4507810]

8: Reports Links
NT_086888

Homo sapiens chromosome 18 genomic contig, alternate assembly
gi|51474639|ref|NT_086888.1|Hs18_86553[51474639]

9: Reports Links
NT_086754

Homo sapiens chromosome 9 genomic contig, alternate assembly
gi|51467694|ref|NT_086754.1|Hs9_86435[51467694]

10: Reports Links
NT_008470

Homo sapiens chromosome 9 genomic contig
gi|51467683|ref|NT_008470.17|Hs9_8627[51467683]

11: Reports Links
NT_010966

Homo sapiens chromosome 18 genomic contig
gi|29823167|ref|NT_010966.13|Hs18_11123[29823167]

12: Reports Links
NC_003071

Arabidopsis thaliana chromosome 2, complete sequence
gi|30698031|ref|NC_003071.3|[30698031]

13: Reports Links
NC_003284

Caenorhabditis elegans chromosome X, complete sequence
gi|32566600|ref|NC_003284.3|[32566600]

14: Reports Links
AJ224156

Rattus norvegicus mRNA for ceramide glucosyltransferase
gi|2924339|emb|AJ224156.1|RNAJ4156[2924339]

15: Reports Links
AF367245

Gossypium arboreum ceramide glucosyltransferase mRNA, complete cds
gi|14718994|gb|AF367245.1|AF367245[14718994]

16: Reports [Links](#)
AF364403
Pichia pastoris ceramide glucosyltransferase gene, complete cds
gi|14718992|gb|AF364403.1|AF364403[14718992]

17: Reports [Links](#)
AF364402
Magnaporthe grisea ceramide glucosyltransferase mRNA, complete cds
gi|14718990|gb|AF364402.1|AF364402[14718990]

18: Reports [Links](#)
AF364401
Caenorhabditis elegans ceramide glucosyltransferase mRNA, complete cds
gi|14718988|gb|AF364401.1|AF364401[14718988]

19: Reports [Links](#)
AF119416
Mus musculus GM3 synthase mRNA, complete cds
gi|7670072|gb|AF119416.1|AF119416[7670072]

20: Reports [Links](#)
AF119415
Homo sapiens GM3 synthase mRNA, complete cds
gi|7670070|gb|AF119415.1|AF119415[7670070]

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Reports

Links

[AF097159](#)

Homo sapiens UDP-Gal:glucosylceramide beta-1,4-galactosyltransferase mRNA, complete cds
gi|5305556|gb|AF097159.1|AF097159[5305556]

 22:

Reports

Links

Related resources

[AF047707](#)

Rattus norvegicus UDP-glucose:ceramide glycosyltransferase mRNA, complete cds
gi|4105566|gb|AF047707.1|AF047707[4105566]

 23:

Reports

Links

[AF097158](#)

Mus musculus UDP-Gal:glucosylceramide beta-1,4-galactosyltransferase mRNA, complete cds
gi|5305554|gb|AF097158.1|AF097158[5305554]

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